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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/544,517

DATE: 03/14/2002

TIME: 14:13:00

Input Set : A:\410.app

Output Set: N:\CRF3\03142002\I544517.raw

4 <110> APPLICANT: Luche, Ralf M.
 5 Wei, Bo
 7 <120> TITLE OF INVENTION: DSP-4 DUAL-SPECIFICITY PHOSPHATASE
 10 <130> FILE REFERENCE: 200125.410
 12 <140> CURRENT APPLICATION NUMBER: US 09/544,517
 13 <141> CURRENT FILING DATE: 2000-04-06
 15 <160> NUMBER OF SEQ ID NOS: 21
 17 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 1087
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Homo sapien
 24 <400> SEQUENCE: 1

25	acattgc	atc	cctggg	ataa	acggac	ctgg	acaact	ctctt	gtgt	gtct	gtct	60
26	cggttac	ctg	gatggg	cgaa	cacctc	tggag	gctggc	ttacc	tggg	aataa	gggac	120
27	tagcagt	tca	gccgtt	ttct	atgcct	gtctg	gatttg	ttttg	ttcc	cagcc	actgc	180
28	tcatgta	atg	tactcc	ctta	accagg	aaat	taaagc	attc	tcccg	gaata	atctc	240
29	gcaatgc	cacc	aggggt	gacaa	cgcta	actgg	aaagaa	aatt	ataga	aacat	ggaa	300
30	cagaatt	cat	gttgtg	gaag	aagtag	agcc	gagcag	tggg	ggtg	gttg	gtgca	360
31	ggacct	tagc	tggac	ctgc	aagttg	gcgt	tatta	agcca	tgg	ttgct	c	420
32	agatgct	gct	catgat	ttgg	atacact	gaa	aaaga	ataag	gtgac	tata	ttct	480
33	tgcatat	gga	gttgaa	aatg	ctttcc	tcag	tgactt	taca	tataa	gagca	ttct	540
34	ggatctg	cct	gaaacca	aaca	tcctgt	ctta	ttttcc	agaa	tg	tttg	aat	600
35	agcaaaa	aga	aaagat	ggag	tggtt	cttgt	tcattg	t	gcagg	cggtt	c	660
36	tgcaatt	gta	ataggt	ttcc	tgatga	attc	tgaaca	aaacc	tcattt	acca	gtgct	720
37	tttggtg	aaa	aatgca	agac	cttccat	atg	tc	caaatt	ct	ggctt	catg	780
38	tacatat	caa	gagggc	aaag	aaagca	ataa	gtgtg	acaga	atac	aggga	acagt	840
39	agttgc	attg	tagcag	acaa	tgga	caactg	tagttt	ctga	attg	actt	ct	900
40	tttccct	ttt	ttggag	agta	gactag	caaa	actcc	cttt	ttct	cttg	cc	960
41	ataaat	ggag	gtcaat	ctga	ttgtc	ctgac	ctact	gtata	agtaa	atttc	aaatg	1020
42	actttc	tctt	tg	tattata	atgtg	tgatt	aaatg	ctttt	ttaa	attg	ct	1080
43	aaaaaaa											1087

45 <210> SEQ ID NO: 2
 46 <211> LENGTH: 217
 47 <212> TYPE: PRT
 48 <213> ORGANISM: Homo sapien
 50 <400> SEQUENCE: 2

51	Met	Tyr	Ser	Leu	Asn	Gln	Glu	Ile	Lys	Ala	Phe	Ser	Arg	Asn	Asn	Leu
52	1				5				10					15		
53	Arg	Lys	Gln	Cys	Thr	Arg	Val	Thr	Thr	Leu	Thr	Gly	Lys	Lys	Ile	Ile
54				20					25					30		
55	Glu	Thr	Trp	Lys	Asp	Ala	Arg	Ile	His	Val	Val	Glu	Glu	Val	Glu	Pro
56				35				40					45			

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57 Ser Ser Gly Gly Gly Cys Gly Tyr Val Gln Asp Leu Ser Ser Asp Leu
58      50                      55                      60
59 Gln Val Gly Val Ile Lys Pro Trp Leu Leu Leu Gly Ser Gln Asp Ala
60 65                      70                      75                      80
61 Ala His Asp Leu Asp Thr Leu Lys Lys Asn Lys Val Thr His Ile Leu
62                      85                      90                      95
63 Asn Val Ala Tyr Gly Val Glu Asn Ala Phe Leu Ser Asp Phe Thr Tyr
64                      100                     105                     110
65 Lys Ser Ile Ser Ile Leu Asp Leu Pro Glu Thr Asn Ile Leu Ser Tyr
66                      115                     120                     125
67 Phe Pro Glu Cys Phe Glu Phe Ile Glu Glu Ala Lys Arg Lys Asp Gly
68                      130                     135                     140
69 Val Val Leu Val His Cys Asn Ala Gly Val Ser Arg Ala Ala Ala Ile
70 145                      150                     155                     160
71 Val Ile Gly Phe Leu Met Asn Ser Glu Gln Thr Ser Phe Thr Ser Ala
72                      165                     170                     175
73 Phe Ser Leu Val Lys Asn Ala Arg Pro Ser Ile Cys Pro Asn Ser Gly
74                      180                     185                     190
75 Phe Met Glu Gln Leu Arg Thr Tyr Gln Glu Gly Lys Glu Ser Asn Lys
76                      195                     200                     205
77 Cys Asp Arg Ile Gln Glu Asn Ser Ser
78      210                      215

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80 <210> SEQ ID NO: 3

81 <211> LENGTH: 14

82 <212> TYPE: PRT

83 <213> ORGANISM: Homo sapien

85 <400> SEQUENCE: 3

86 Val His Cys Asn Ala Gly Val Ser Arg Ala Ala Ala Ile Val

87 1 5 10

89 <210> SEQ ID NO: 4

90 <211> LENGTH: 23

91 <212> TYPE: PRT

92 <213> ORGANISM: Homo sapien

94 <400> SEQUENCE: 4

95 Gly Arg Val Leu Val His Cys Gln Ala Gly Ile Ser Arg Ser Gly Thr

96 1 5 10 15

97 Asn Ile Leu Ala Tyr Leu Met

98 20

100 <210> SEQ ID NO: 5

101 <211> LENGTH: 27

102 <212> TYPE: DNA

103 <213> ORGANISM: Homo sapien

105 <400> SEQUENCE: 5

106 gcataaaaaa ggcaagagaa aaaaggg

108 <210> SEQ ID NO: 6

109 <211> LENGTH: 24

110 <212> TYPE: DNA

111 <213> ORGANISM: Homo sapien

113 <400> SEQUENCE: 6

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114	cacacttatt gctttctttg ccct	24
116	<210> SEQ ID NO: 7	
117	<211> LENGTH: 25	
118	<212> TYPE: DNA	
119	<213> ORGANISM: Homo sapien	
121	<400> SEQUENCE: 7	
122	gcactggtaa atgaggtttg ttcag	25
124	<210> SEQ ID NO: 8	
125	<211> LENGTH: 18	
126	<212> TYPE: DNA	
127	<213> ORGANISM: Homo sapien	
129	<400> SEQUENCE: 8	
130	agccctggaa acgcctgc	18
132	<210> SEQ ID NO: 9	
133	<211> LENGTH: 25	
134	<212> TYPE: DNA	
135	<213> ORGANISM: Homo sapien	
137	<400> SEQUENCE: 9	
138	ctgaacaaac ctcatttacc agtgc	25
140	<210> SEQ ID NO: 10	
141	<211> LENGTH: 18	
142	<212> TYPE: DNA	
143	<213> ORGANISM: Homo sapien	
145	<400> SEQUENCE: 10	
146	gcaggcgttt ccagggt	18
148	<210> SEQ ID NO: 11	
149	<211> LENGTH: 25	
150	<212> TYPE: DNA	
151	<213> ORGANISM: Homo sapien	
153	<400> SEQUENCE: 11	
154	gagggcacaag aaagcaataa gtgtg	25
156	<210> SEQ ID NO: 12	
157	<211> LENGTH: 27	
158	<212> TYPE: DNA	
159	<213> ORGANISM: Homo sapien	
161	<400> SEQUENCE: 12	
162	gtcacactta ttgctttctt tgccctc	27
164	<210> SEQ ID NO: 13	
165	<211> LENGTH: 20	
166	<212> TYPE: DNA	
167	<213> ORGANISM: Homo sapien	
169	<400> SEQUENCE: 13	
170	gcaggcgttt ccagggtgc	20
172	<210> SEQ ID NO: 14	
173	<211> LENGTH: 170	
174	<212> TYPE: PRT	
175	<213> ORGANISM: Homo sapien	
177	<400> SEQUENCE: 14	
178	Ser Asp Leu Asp Arg Asp Pro Asn Ser Ala Thr Asp Ser Asp Gly Ser	

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179 1          5          10          15
180 Pro Leu Ser Asn Ser Gln Pro Ser Phe Pro Val Glu Ile Leu Pro Phe
181          20          25          30
182 Leu Tyr Leu Gly Cys Ala Lys Asp Ser Thr Asn Leu Asp Val Leu Glu
183          35          40          45
184 Glu Phe Gly Ile Lys Tyr Ile Leu Asn Val Thr Pro Asn Leu Pro Asn
185          50          55          60
186 Leu Phe Glu Asn Ala Gly Glu Phe Lys Tyr Lys Gln Ile Pro Ile Ser
187 65          70          75          80
188 Asp His Trp Ser Gln Asn Leu Ser Gln Phe Phe Pro Glu Ala Ile Ser
189          85          90          95
190 Phe Ile Asp Glu Ala Arg Gly Lys Asn Cys Gly Val Leu Val His Cys
191          100          105          110
192 Leu Ala Gly Ile Ser Arg Ser Val Thr Val Thr Val Ala Tyr Leu Met
193          115          120          125
194 Gln Lys Leu Asn Leu Ser Met Asn Asp Ala Tyr Asp Ile Val Lys Met
195          130          135          140
196 Lys Lys Ser Asn Ile Ser Pro Asn Phe Asn Phe Met Gly Gln Leu Leu
197 145          150          155          160
198 Asp Phe Glu Arg Thr Leu Gly Leu Ser Ser
199          165          170
201 <210> SEQ ID NO: 15
202 <211> LENGTH: 168
203 <212> TYPE: PRT
204 <213> ORGANISM: Homo sapien
206 <400> SEQUENCE: 15
207 Asp Arg Glu Leu Pro Ser Ser Ala Thr Glu Ser Asp Gly Ser Pro Val
208 1          5          10          15
209 Pro Ser Ser Gln Pro Ala Phe Pro Val Gln Ile Leu Pro Tyr Leu Tyr
210          20          25          30
211 Leu Gly Cys Ala Lys Asp Ser Thr Asn Leu Asp Val Leu Gly Lys Tyr
212          35          40          45
213 Gly Ile Lys Tyr Ile Leu Asn Val Thr Pro Asn Leu Pro Asn Ala Phe
214          50          55          60
215 Glu His Gly Gly Glu Phe Thr Tyr Lys Gln Ile Pro Ile Ser Asp His
216 65          70          75          80
217 Trp Ser Gln Asn Leu Ser Gln Phe Phe Pro Glu Ala Ile Ser Phe Ile
218          85          90          95
219 Asp Glu Ala Arg Ser Lys Lys Cys Gly Val Leu Val His Cys Leu Ala
220          100          105          110
221 Gly Ile Ser Arg Ser Val Thr Val Thr Val Ala Tyr Leu Met Gln Lys
222          115          120          125
223 Met Asn Leu Ser Leu Asn Asp Ala Tyr Asp Phe Val Lys Arg Lys Lys
224          130          135          140
225 Ser Asn Ile Ser Pro Asn Phe Asn Phe Met Gly Gln Leu Leu Asp Phe
226 145          150          155          160
227 Glu Arg Thr Leu Gly Leu Ser Ser
228          165
230 <210> SEQ ID NO: 16

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231 <211> LENGTH: 170
232 <212> TYPE: PRT
233 <213> ORGANISM: Homo sapien
235 <400> SEQUENCE: 16
236 Gly Leu Cys Glu Gly Lys Pro Ala Ala Leu Leu Pro Met Ser Leu Ser
237 1 5 10 15
238 Gln Pro Cys Leu Pro Val Pro Ser Val Gly Leu Thr Arg Ile Leu Pro
239 20 25 30
240 His Leu Tyr Leu Gly Ser Gln Lys Asp Val Leu Asn Lys Asp Leu Met
241 35 40 45
242 Thr Gln Asn Gly Ile Ser Tyr Val Leu Asn Ala Ser Asn Ser Cys Pro
243 50 55 60
244 Lys Pro Asp Phe Ile Cys Glu Ser Arg Phe Met Arg Val Pro Ile Asn
245 65 70 75 80
246 Asp Asn Tyr Cys Glu Lys Leu Leu Pro Trp Leu Asp Lys Ser Ile Glu
247 85 90 95
248 Phe Ile Asp Lys Ala Lys Leu Ser Ser Cys Gln Val Ile Val His Cys
249 100 105 110
250 Leu Ala Gly Ile Ser Arg Ser Ala Thr Ile Ala Ile Ala Tyr Ile Met
251 115 120 125
252 Lys Thr Met Gly Met Ser Ser Asp Asp Ala Tyr Arg Phe Val Lys Asp
253 130 135 140
254 Arg Arg Pro Ser Ile Ser Pro Asn Phe Asn Phe Leu Gly Gln Leu Leu
255 145 150 155 160
256 Glu Tyr Glu Arg Thr Leu Lys Leu Leu Ala
257 165 170
259 <210> SEQ ID NO: 17
260 <211> LENGTH: 168
261 <212> TYPE: PRT
262 <213> ORGANISM: Homo sapien
264 <400> SEQUENCE: 17
265 Pro Ala Gln Ala Leu Pro Pro Ala Gly Ala Glu Asn Ser Asn Ser Asp
266 1 5 10 15
267 Pro Arg Val Pro Ile Tyr Asp Gln Gly Gly Pro Val Glu Ile Leu Pro
268 20 25 30
269 Tyr Leu Tyr Leu Gly Ser Cys Asn His Ser Ser Asp Leu Gln Gly Leu
270 35 40 45
271 Gln Ala Cys Gly Ile Thr Ala Val Leu Asn Val Ser Ala Ser Cys Pro
272 50 55 60
273 Asn His Phe Glu Gly Leu Phe His Tyr Lys Ser Ile Pro Val Glu Asp
274 65 70 75 80
275 Asn Gln Met Val Glu Ile Ser Ala Trp Phe Gln Glu Ala Ile Ser Phe
276 85 90 95
277 Ile Asp Ser Val Lys Asn Ser Gly Gly Arg Val Leu Val His Cys Gln
278 100 105 110
279 Ala Gly Ile Ser Arg Ser Ala Thr Ile Cys Leu Ala Tyr Leu Ile Gln
280 115 120 125
281 Ser His Arg Val Arg Leu Asp Glu Ala Phe Asp Phe Val Lys Gln Arg
282 130 135 140

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VERIFICATION SUMMARY

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